

## SEQUENCE LISTING

<110> Lambeth, J. David  
Cheng, Guangjie

<120> Regulatory Protein For Nox Enzymes

<130> 05501-0202 (43150-287577)

<150> US 60/405,647

<151> 2002-08-23

<150> US 60/396,170

<151> 2002-07-16

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<170> PatentIn version 3.1

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225 230 235	
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Gln Phe Cys Ala Ser Arg Ala Tyr Glu Ser Ser Arg Ala Asp Glu Leu	
240 245 250 255	
tcc gtg ccc gcg ggg gcg cgc gtg cgc gtg ttg gaa acg tca gac cgc	817
Ser Val Pro Ala Gly Ala Arg Val Arg Val Leu Glu Thr Ser Asp Arg	
260 265 270	
ggc tgg tgg cta tgc agg tac ggc gac cgg gcg ggc cta ctc ccc gcg	865
Gly Trp Trp Leu Cys Arg Tyr Gly Asp Arg Ala Gly Leu Leu Pro Ala	
275 280 285	
gtg ctg ctg cgg ccg gaa ggg ctg ggc gct ctc ctg agc ggg acg ggg	913

Val	Leu	Leu	Arg	Pro	Glu	Gly	Leu	Gly	Ala	Leu	Leu	Ser	Gly	Thr	Gly		
		290					295					300					
ttc	cgt	gga	gga	gac	gac	ccg	gcg	ggg	gag	gcc	cgg	ggc	ttc	cct	gaa		961
Phe	Arg	Gly	Gly	Asp	Asp	Pro	Ala	Gly	Glu	Ala	Arg	Gly	Phe	Pro	Glu		
	305					310					315						
ccc	tcc	cag	gcc	acc	gcc	cct	ccc	ccc	acc	gtg	ccc	acc	cga	cct	tcg		1009
Pro	Ser	Gln	Ala	Thr	Ala	Pro	Pro	Pro	Thr	Val	Pro	Thr	Arg	Pro	Ser		
320					325					330					335		
ccg	ggc	gcc	atc	cag	agc	cgc	tgc	tgc	acc	gtc	aca	cgc	agg	gcc	ctg		1057
Pro	Gly	Ala	Ile	Gln	Ser	Arg	Cys	Cys	Thr	Val	Thr	Arg	Arg	Ala	Leu		
				340					345					350			
gag	cgg	cgc	cca	cgg	cgc	cag	ggc	cgc	cct	cga	ggg	tgc	gtg	gac	tct		1105
Glu	Arg	Arg	Pro	Arg	Arg	Gln	Gly	Arg	Pro	Arg	Gly	Cys	Val	Asp	Ser		
			355					360					365				
gtg	ccg	cac	ccc	acg	acg	gag	cag	tgagcgcgag	gatcc								1144
Val	Pro	His	Pro	Thr	Thr	Glu	Gln										
		370				375											

&lt;210&gt; 8

&lt;211&gt; 375

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 8

Met	Ala	Gly	Pro	Arg	Tyr	Pro	Val	Ser	Val	Gln	Gly	Ala	Ala	Leu	Val		
1				5				10						15			
Gln	Ile	Lys	Arg	Leu	Gln	Thr	Phe	Ala	Phe	Ser	Val	Arg	Trp	Ser	Asp		
		20						25					30				
Gly	Ser	Asp	Thr	Phe	Val	Arg	Arg	Ser	Trp	Asp	Glu	Phe	Arg	Gln	Leu		
		35					40					45					
Lys	Thr	Leu	Lys	Glu	Thr	Phe	Pro	Val	Glu	Ala	Gly	Leu	Leu	Arg	Arg		
	50					55					60						
Ser	Asp	Arg	Val	Leu	Pro	Lys	Leu	Leu	Gly	Gln	Ala	Ser	Leu	Asp	Ala		
65					70					75					80		



Pro Leu Leu Gly Arg Val Gly Arg Thr Ser Arg Gly Leu Ala Arg Leu  
85 90 95

Gln Leu Leu Glu Thr Tyr Ser Arg Arg Leu Leu Ala Thr Ala Glu Arg  
100 105 110

Val Ala Arg Ser Pro Thr Ile Thr Gly Phe Phe Ala Pro Gln Pro Leu  
115 120 125

Asp Leu Glu Pro Ala Leu Pro Pro Gly Ser Arg Val Ile Leu Pro Thr  
130 135 140

Pro Glu Glu Gln Pro Leu Ser Arg Ala Ala Gly Arg Leu Ser Ile His  
145 150 155 160

Ser Leu Glu Ala Gln Ser Leu Arg Cys Leu Gln Pro Phe Cys Thr Gln  
165 170 175

Asp Thr Arg Asp Arg Pro Phe Gln Ala Gln Ala Gln Glu Ser Leu Asp  
180 185 190

Val Leu Leu Arg His Pro Ser Gly Trp Trp Leu Val Glu Asn Glu Asp  
195 200 205

Arg Gln Thr Ala Trp Phe Pro Ala Pro Tyr Leu Glu Glu Ala Ala Pro  
210 215 220

Gly Gln Gly Arg Glu Gly Gly Pro Ser Leu Gly Ser Ser Gly Pro Gln  
225 230 235 240

Phe Cys Ala Ser Arg Ala Tyr Glu Ser Ser Arg Ala Asp Glu Leu Ser  
245 250 255

Val Pro Ala Gly Ala Arg Val Arg Val Leu Glu Thr Ser Asp Arg Gly  
260 265 270

Trp Trp Leu Cys Arg Tyr Gly Asp Arg Ala Gly Leu Leu Pro Ala Val  
275 280 285

Leu Leu Arg Pro Glu Gly Leu Gly Ala Leu Leu Ser Gly Thr Gly Phe  
 290 295 300

Arg Gly Gly Asp Asp Pro Ala Gly Glu Ala Arg Gly Phe Pro Glu Pro  
 305 310 315 320

Ser Gln Ala Thr Ala Pro Pro Pro Thr Val Pro Thr Arg Pro Ser Pro  
 325 330 335

Gly Ala Ile Gln Ser Arg Cys Cys Thr Val Thr Arg Arg Ala Leu Glu  
 340 345 350

Arg Arg Pro Arg Arg Gln Gly Arg Pro Arg Gly Cys Val Asp Ser Val  
 355 360 365

Pro His Pro Thr Thr Glu Gln  
 370 375

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24

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<212> DNA

<213> Artificial Sequence

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<223> Synthetic primer

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gtccatcccc tcatcgggat cctc

24

<210> 11

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic primer

<400> 11

tcaggaatct gcagcctgga agcc

24